

- 1 -

piece 1, NC_000913, yhcC_gltB-, config: linear, direction: -, begin: 3352683, end: 3352053

...] NC_000913.gltB

p35 4.1 bits

... p10

1

1

--> p35-(22)-p10 3352604 Gap 2.3 bits
--> p35-p10 3352604 total 4.3 bits

The diagram illustrates the 3352600 mRNA sequence with its 5' and 3' ends. The sequence is color-coded by codon: green for gln, blue for asp, red for lys, purple for arg, pink for ser, orange for leu, yellow for tyr, light blue for his, and grey for pro. Asterisks indicate start sites at positions 3352590, 3352580, 3352570, 3352560, 3352550, 3352540, and 3352530. A red circle highlights the first start site at position 3352590. Below the sequence, the corresponding amino acid sequence is shown: -val-lys-arg-gly-ser-gln-leu-leu-his-lys-tyr-asp-glutamyl-gln-alanine-alanine-his-pro-glutamyl-ser-phe-gln-argin-ile-ser-asparagine-leu-. The start site at position 3352590 is preceded by a fMet residue.

... + p10 2.5 bits

UW-Madison p10

... p35

ANSWER The answer is 1000. The area of the rectangle is $10 \times 100 = 1000$.

p35-p10 3352460 total 5.3 bits(-----) ... p35-(23)-p10 3352421 Gap
p35 1.4 bits |----- ... p35-p10 3352421 total 6.4

{ p35-(23)-p10 3352433 Gap
p35-p10 3352433 total 7.5

Sequence logo showing the codon context for the first amino acid of each protein. The sequence starts with fMet, followed by various amino acids. The logo shows the frequency of each nucleotide at each position relative to the start site.

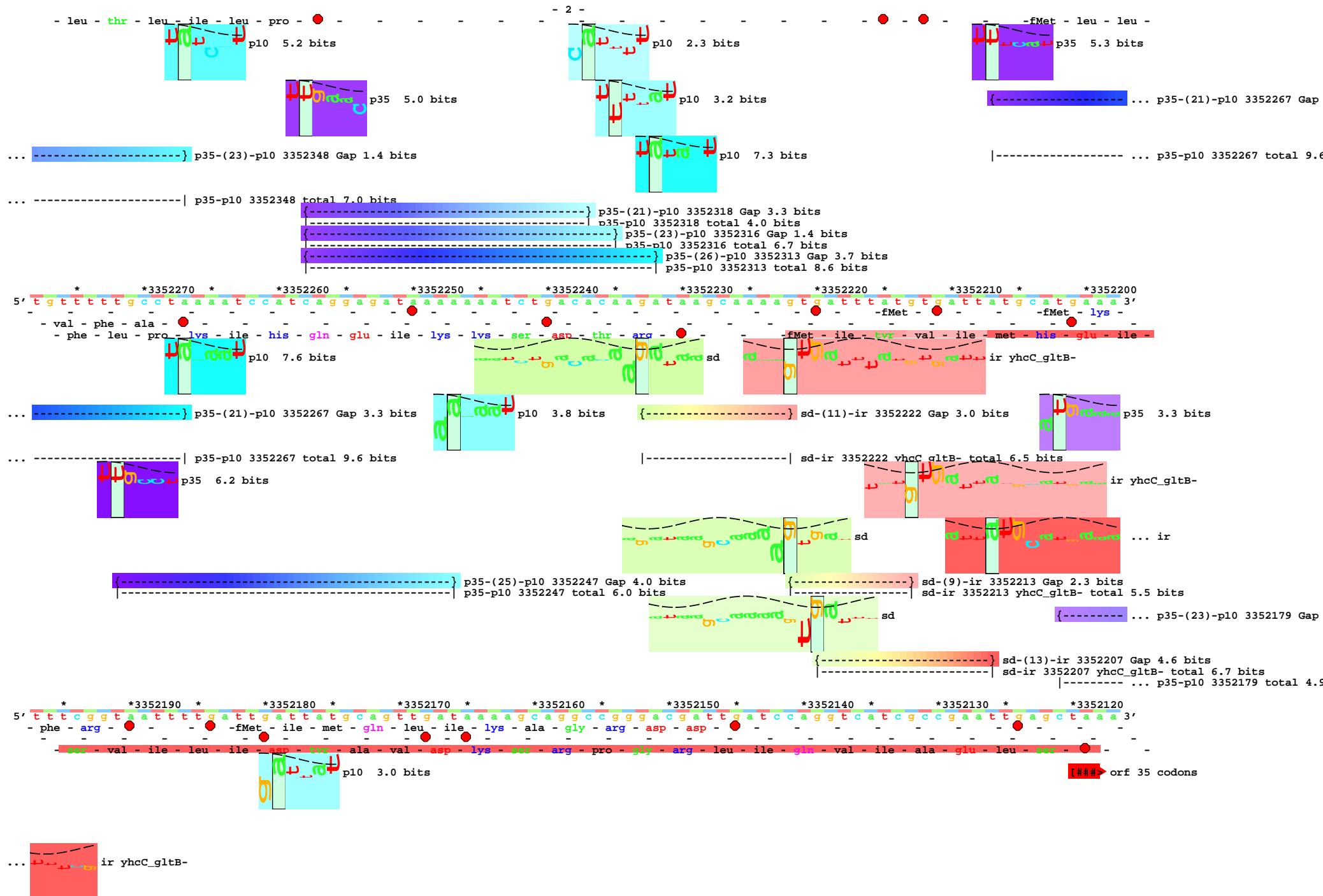
...  yhcC_gltB- [###> orf 19 codons

... p35 3.4 bits p10 4.5 bits p35 3.2 bits

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{-----} sd-(9)-ir 3352430 Gap 2.3 bits  
{-----} p25-(23)-p10 3352431 Gap 1.4 bits
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p35-(23)-p10 3352433 Gap 1.4 bits
p35-p10 3352421 total 6.4 bits
p35-(23)-p10 3352433 Gap 1.4 bits
p35-p10 3352433 total 7.5 bits

* \$132552350 * \$132552310 * \$132552320 * \$132552330 * \$132552310 * \$132552320 * \$132552330 * \$132552340



... -----} p35-(23)-p10 3352179 Gap 1.4 bits

... -----| p35-p10 3352179 total 4.9 bits

* *3352110 * *3352100 * *3352090 * *3352080 * *3352070 * *3352060 *
5' a a t g g c a g g c t t t g g c c t t t t c a a g g c g g g t c t a t c a g a t t a t g c a g t t a c a g a a a t t a g t 3'
-fMet - ala - gly - phe - gly - leu - phe - ser - arg - arg - gly - leu - ser - asp - tyr - ala - val - thr - glu - ile - ser -
- - - -fMet - ala - ser - phe - gln - gly - ala - val - tyr - gln - ile - met - gln - leu - gln - lys - leu -

p35 4.6 bits

[----- ... NC_000913.yhcC

p10 3.0 bits

{-----} p35-(21)-p10 3352075 Gap 3.3 bits
p35-p10 3352075 total 4.4 bits